

SEQUENCE LISTING

<110> Vaisvila, Romualdus Morgan, Richard D. Kucera, Rebecca B. Claus, Toby B. Raleigh, Elisabeth A. Method For Cloning And Producing The MseI Restriction <120> Endonuclease NEB-181 <130> <140> US 09/689,343 <141> 2000-10-12 <160> 21 <170> PatentIn version 3.1 <210> 1 <211> 903 <212> DNA <213> Micrococcus sp. <220> <221> CDS <222> (1)..(900)<223> <400> 1 atg cct atc tcg acc gtc tgg acg ccg gac gga gac gac ctc atc gtg 48 Met Pro Ile Ser Thr Val Trp Thr Pro Asp Gly Asp Asp Leu Ile Val gag gcg gac aac ctc gat ttc att caa acg ctc ccc gac gcg agc ttc 96 Glu Ala Asp Asn Leu Asp Phe Ile Gln Thr Leu Pro Asp Ala Ser Phe 20 25 30 cga atg atc tac atc gat ccg ccg ttc aac aca ggg cga acg cag cgg 144 Arg Met Ile Tyr Ile Asp Pro Pro Phe Asn Thr Gly Arg Thr Gln Arg 40 192 ctt cag tcg ctc aag acg acc cgc tcg gtc aca ggg tcg cga gtc ggc Leu Gln Ser Leu Lys Thr Thr Arg Ser Val Thr Gly Ser Arg Val Gly ttc aaa ggc cag acg tac gac acg gtc aag agc act ctg cac tcg tat 240 Phe Lys Gly Gln Thr Tyr Asp Thr Val Lys Ser Thr Leu His Ser Tyr 65 gac gac gct ttc acc gac tat tgg tcg ttc ctc gaa ccg cgt ctc ctg 288

Asp Asp Ala Phe Thr Asp Tyr Trp Ser Phe Leu Glu Pro Arg Leu Leu

90

95

85



andt t

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gcg cgc tcg Ala Arg Ser 145	Lys Ser Ly			480
tat gtg aag Tyr Val Lys				528
cgc gag ccc Arg Glu Pro				576
ctt ggc aag Leu Gly Lys 195	Leu Pro T	Trp Trp		624
gcg agc aaa Ala Ser Lys 210				672
atc cgt cgc Ile Arg Arg 225	g Met Ile G			720
gat ttc ttc Asp Phe Phe				768
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Arg Met Ile Tyr Ile Asp Pro Pro Phe Asn Thr Gly Arg Thr Gln Arg 35 40 45

Leu Gln Ser Leu Lys Thr Thr Arg Ser Val Thr Gly Ser Arg Val Gly 50 55 60

Phe Lys Gly Gln Thr Tyr Asp Thr Val Lys Ser Thr Leu His Ser Tyr 65 70 75 80

Asp Asp Ala Phe Thr Asp Tyr Trp Ser Phe Leu Glu Pro Arg Leu Leu 85 90 95

Glu Ala Trp Arg Leu Leu Thr Pro Asp Gly Ala Leu Tyr Leu His Leu
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Asp Tyr Arg Glu Val His Tyr Ala Lys Val Val Leu Asp Ala Met Phe 115 120 125

Gly Arg Glu Ser Phe Leu Asn Glu Leu Ile Trp Ala Tyr Asp Tyr Gly
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Ala Arg Ser Lys Ser Lys Trp Pro Thr Lys His Asp Asn Ile Leu Val 145 150 155 160

Tyr Val Lys Asp Pro Asn Asn Tyr Val Trp Asn Gly Gln Asp Val Asp 165 170 175

Arg Glu Pro Tyr Met Ala Pro Gly Leu Val Thr Pro Glu Lys Val Ala 180 185 190

Leu Gly Lys Leu Pro Thr Asp Val Trp Trp His Thr Ile Val Pro Pro 195 200 205

Ala Ser Lys Glu Arg Thr Gly Tyr Ala Thr Gln Lys Pro Val Gly Ile 210 215 220

Ile Arg Arg Met Ile Gln Ala Ser Ser Asn Glu Gly Asp Trp Val Leu 225 230 235 240

Asp Phe Phe Ala Gly Ser Gly Thr Thr Gly Ala Ala Ala Arg Gln Leu 245 . 250 . 255

Gly Arg Arg Phe Val Leu Val Asp Val Asn Pro Glu Ala Ile Ala Val 260 265 270

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80

tcg cgt cgt cca caa acc tgg cag ctt gca gaa gga tat acg gac gag

Ser Arg Arg Pro Gln Thr Trp Gln Leu Ala Glu Gly Tyr Thr Asp Glu

85 90 95

Phe Phe Thr Asp Arg Thr Tyr Ala Ala Arg Ile Gly His Gly Glu Asp

70

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Trp Lys Asp Leu Asp Glu Tyr Leu Asp Phe Leu Tyr Pro Arg Leu Val
100 105 110

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						ctc Leu 150											480
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				_	_	acn Xaa											624
·						ccg Pro											672
					_	gtc Val 230											720
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		_		_		gat Asp	_		-								816
						gca Ala											864
						gcc Ala											912
						act Thr 310											960
						cca Pro											1008
	cgc	ctc	caa	gca	ccc	ttt	ctc	gta	gat	ttt	tgg	gaa	gtg	gac	gat	caa	1056

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Arg Leu Gln Ala Pro Phe Leu Val Asp Phe Trp Glu Val Asp Asp Gln 345 340 1104 tgg gat ggc aaa atc ttc cgc agc cgt cat caa ggc tta cgc tcc cgc Trp Asp Gly Lys Ile Phe Arg Ser Arg His Gln Gly Leu Arg Ser Arg 360 355 1152 ctt cag gag cag gcg ccg ctc tct cta cca ttg acc ggg aat gga ctg Leu Gln Glu Gln Ala Pro Leu Ser Leu Pro Leu Thr Gly Asn Gly Leu 375 1200 ttg tgt gta cgg gta gtg agc cgt gaa ggg gaa tac tat gag ttc aca Leu Cys Val Arg Val Val Ser Arg Glu Gly Glu Tyr Tyr Glu Phe Thr 395 390 1236 ggt cga gcc gat agc cct cac ccc gta tcg ttt tga Gly Arg Ala Asp Ser Pro His Pro Val Ser Phe 405 410 <210> 4 <211> 411 <212> PRT <213> Unknown <220> <223> Environmental DNA <220> <221> misc_feature <222> (198)..(198) <223> Xaa = any amino acid <400> 4 Met Pro Thr Leu Asp Trp Pro Gly Lys Gln Leu Ser Phe Pro Pro Ala Thr Ser Leu His Leu Glu Ser Val Val Thr Glu Gly Ala Glu Ser Pro 25 30 Pro Asn Arg Leu Ile Trp Ala Asp Asn Leu Pro Leu Met Val Asp Leu 40 Leu Ala Glu Tyr Glu Gly Lys Ile Asp Leu Ile Tyr Ala Asp Pro Pro 55 Phe Phe Thr Asp Arg Thr Tyr Ala Ala Arg Ile Gly His Gly Glu Asp 70 Ser Arg Arg Pro Gln Thr Trp Gln Leu Ala Glu Gly Tyr Thr Asp Glu Trp Lys Asp Leu Asp Glu Tyr Leu Asp Phe Leu Tyr Pro Arg Leu Val 105 110

Leu Met Tyr Arg Leu Leu Ala Pro His Gly Thr Leu Tyr Leu His Leu

115 120 125

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Gly 145	Arg	Gln	Arg	Phe	Leu 150	Asn	Glu	Ile	Val	Trp 155	Ile	Tyr	His	Gly	Pro 160
Ser	Ala	Ile	Arg	Arg 165	Ala	Phe	Lys	Arg	Lys 170	His	Asp	Thr	Ile	Leu 175	Val
Tyr	Val	Lys	Gly 180	Glu	Asn	Tyr	Thr	Phe 185	Asn	Ala	Asp	Ala	Val 190	Arg	Glr
Pro	Tyr	His 195	Pro	Ser	Xaa	His	Lys 200	Thr	Phe	Ala	Ser	Ser 205	Pro	Lys	Ala
Gly	Phe 210	Gly	Lys	Val	Pro	Asp 215	Leu	Gln	Arg	Gly	Lys 220	Val	Pro	Glu	Asp
Trp 225	Trp	Tyr	Phe	Pro	Val 230	Val	Ala	Arg	Leu	His 235	Arg	Glu	Arg	Ser	Gly 240
Tyr	Pro	Thr	Gln	Lys 245	Pro	Gln	Ala	Leu	Leu 250	Glu	Arg	Ile	Leu	Leu 255	Ala
Ser	Ser	Asn	Ala 260	Gly	Asp	Leu	Val	Ala 265	Asp	Phe	Phe	Cys	Gly 270	Ser	Gly
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Glu 305	Gly	Val	Ser	Phe	Thr 310	Phe	Glu	Arg	Gln	Glu 315	Thr	Phe	Thr	Leu	Pro 320
Ile	Gln	Pro	Leu	Pro 325	Pro	Asp	Trp	Leu	Ile 330	Ile	Ala	Glu	Glu	Gln 335	Il∈
Arg	Leu	Gln	Ala 340	Pro	Phe	Leu	Val	Asp 345	Phe	Trp	Glu	Val	Asp 350	Asp	Glr
Trp	Asp	Gly 355	Lys	Ile	Phe	Arg	Ser 360	Arg	His	Gln	Gly	Leu 365	Arg	Ser	Arg
Leu	Gln 370	Glu	Gln	Ala	Pro	Leu 375	Ser	Leu	Pro	Leu	Thr 380	Gly	Asn	Gly	Leu
Leu 385	Cys	Val	Arg	Val	Val 390	Ser	Arg	Glu	Gly	Glu 395	Tyr	Tyr	Glu	Phe	Thr
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Ala Asp Asn Met Glu Val Leu Arg Gly Leu Pro Ala Ala Ser Val Asp
ctg atc tac atc gat cct ccg ttc aac acc gga aag gtt cag gag cgc
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Leu Ile Tyr Ile Asp Pro Pro Phe Asn Thr Gly Lys Val Gln Glu Arg
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act cag ctc aaa acg gtg cgc tcc gag tgg ggc gat cgc gtc gga ttc
                                                                       192
Thr Gln Leu Lys Thr Val Arg Ser Glu Trp Gly Asp Arg Val Gly Phe
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                        55
cag ggc cgt cgc tac gaa agc atc gtc gtg ggt aag aag cgc ttt acc
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Gln Gly Arg Arg Tyr Glu Ser Ile Val Val Gly Lys Lys Arg Phe Thr
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gac ttc ttc gac gac tat ctg gct ttc ctg gaa ccg cgc ctg gtc gaa
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Asp Phe Phe Asp Asp Tyr Leu Ala Phe Leu Glu Pro Arg Leu Val Glu
                85
                                                         95
                                                                       336
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Ala His Arg Val Leu Ala Pro His Gly Cys Leu Tyr Phe His Val Asp
            100
                                                                       384
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Tyr Arg Glu Val His Tyr Cys Lys Val Leu Leu Asp Gly Ile Phe Gly
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                            120
cgc gag gcc ttt ctc aac gag atc atc tgg gcc tac gat tac ggc ggg
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Arg Glu Ala Phe Leu Asn Glu Ile Ile Trp Ala Tyr Asp Tyr Gly Gly
                        135
cgt ccg aag gac agg tgg cct cct aag cac gac aac atc ctg ctc tac
                                                                       480
Arg Pro Lys Asp Arg Trp Pro Pro Lys His Asp Asn Ile Leu Leu Tyr
145
                    150
                                         155
                                                                       528
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Leu Ile Tyr Ile Asp Pro 35	Pro Phe Asn Thr 40	Gly Lys Val Gln Glu A 45	rg

En

Thr Gln Leu Lys Thr Val Arg Ser Glu Trp Gly Asp Arg Val Gly Phe 50 55 60

Gln Gly Arg Arg Tyr Glu Ser Ile Val Val Gly Lys Lys Arg Phe Thr 65 70 75 80

Asp Phe Phe Asp Asp Tyr Leu Ala Phe Leu Glu Pro Arg Leu Val Glu 85 90 95

Ala His Arg Val Leu Ala Pro His Gly Cys Leu Tyr Phe His Val Asp 100 105 110

Tyr Arg Glu Val His Tyr Cys Lys Val Leu Leu Asp Gly Ile Phe Gly 115 120 125

Arg Glu Ala Phe Leu Asn Glu Ile Ile Trp Ala Tyr Asp Tyr Gly Gly
130 135 140

Arg Pro Lys Asp Arg Trp Pro Pro Lys His Asp Asn Ile Leu Leu Tyr 145 150 155 160

Ala Lys Thr Pro Gly Arg His Val Phe Asn Ala Asp Glu Ile Glu Arg 165 170 175

Ile Pro Tyr Met Ala Pro Gly Leu Val Gly Pro Glu Lys Ala Ala Arg 180 185 190

Gly Lys Leu Pro Thr Asp Thr Trp Trp His Thr Ile Val Pro Thr Ser 195 200 205

Gly Ser Glu Lys Thr Gly Tyr Pro Thr Gln Lys Pro Leu Gly Ile Leu 210 215 220

Arg Arg Ile Val Gln Ala Ser Ser His Pro Gly Ala Val Leu Asp 225 230 235 240

Phe Phe Ala Gly Ser Gly Thr Thr Gly Val Ala Ala Phe Glu Leu Gly 245 250 255

Arg Arg Phe Ile Leu Val Asp Asn His Pro Glu Ala Leu Gln Val Met 260 265 270

Ala Arg Arg Phe Asp Gly Ile Glu Gly Ile Glu Trp Val Gly Phe Asp 275 280 285

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185

180

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